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1 GCGGCCGCGT CGACGTCTTT GCTGCCGCAC AGGGAGCAGC AGCAGCCGCC GACCCGATCC
61 CTTGGGAGCC CACCAAGTGC TCGCCTTGCT TAGCAGCTAC AGGAGCTGCC GCGGGGTTGC
121 TCCCTGAGGC AGCGTGCA TGATGGTCCG CAGCCAGCTT GGTGTGCGAG CCGTACTTCT
181 TGGAAGCGAG AGAGACTGTG GGAGAGCGCA AATCACTCCA GCCGCTTCCA GGGGAGTCTG
241 GGGACCGCAG GAGCGTTGGA GGCTGCCTGC CGGCATAAAC AGGAACAAGC GCATTCTTAT
301 TCTTCTGTGG TTGCTGAGTT CTGGCTGCGT TCAAGGGGGT TCACCTCTTC CCCTTCTGGC
361 GAGTTTTTGC TGCGTCTTTC CCTAAGAAGC AGCGCCACGT GCGTGGCGTG CCTCAGCCTG
421 ACGCGGTGCA CCTTTTACGT AAGAGCGTCG ATAGCATCGG TCATCTACAG CAGCGTGCTG
481 CTGCTTCCGT GACCTTTACA CTGCTTGTTG CGGGCCGTCT TGTAGAGGGG CCATCTGCTT
541 GTTCGCTGCT GGACGCAGAC CCGGCGCCCG ACATTTCCGG CAGCCGGGCA GTTGAGATAA
601 ACCGGCTGCC CGGTGGCCGT CGAAATTGAA GCAGGATCTC TACAGTAAGG AACAAATCGC
661 GCTATTTTTA AGGAGTGTGT ATACTTGGGG CGTTACTCGT GAGTATTGCT GATGATGGAC
721 GTCCGTGTGG GGGGTAAGTA TCGTTTGGGG AGGAAGATTG GGAGCGGATC CTTCGGCGAC
781 ATCTACCTTG GTACGAACAT CTCAACAGGA GATGAAGTCG CTATCAAATT GGAAAGCGTG
841 CGGTCTAGGC ATCCACAAC TATCTATGAA AGCAAGCTGT ACAAATCCT AACGGGTGGA
901 ATCGGAATCC CGACTCTTTA CTGGTATGGG ATCGAGGGGG ATTACAACGT TATGATTATT
961 GAGCTTTTGG GCCCGTCTCT TGAGGACCTC TTCAGCATTT GCAACAGAAA GCTTTCTTTG
1021 AAGACTGTTC TGATGCTCGC CGACCAAATG CTAAATCGTA TTGAGTTCGT CCACAGCAGA
1081 CATTTTCATCC ATCGAGACAT CAAGCCTGAC AATTTTTTGA TCGGTAGGGG CAAAAGATG
1141 TCCATTGTTT TTGCTATCGA CTTTGGCCTC GCAAAGAAGT ACAGAGATCC CAGAACACAG
1201 TCCCATATTC CTTATCGAGA AGGGAAGAAC CTGACAGGTA CCGCGAGGTA CGCCTCTGTG
1261 AACACCCACT TGGGAATAGA ACAGAGCAGG CGCGATGATC TGGAAGCGCT CGGCTACGTC
1321 TTAATGTACT TCAACAGAGG TTCCTTACCC TGGCAAGGAT TAAAGGCCAC TACGAAGAAA
1381 GATAAATATG ACAAGATTAT GGAGAAGAAG ATGTCCACCC CTATTGAAGT CCTTTGCAAA
1441 CAATTTCCAT TTGAGTTTAT CACATATCTG AACTATTGCC GGTCTCTGCG ATTCTGAAGAT
1501 CGCCCGGACT ATTCTTATTT GAGACGGTTG TTCAAGGATC TTTTCTTCCG TGAGGGATAC
1561 CAGTATGACT TTATATTCGA TTGGACATTT CTGCATGCTG AGAGAGAGCG CGAGCGTCAA
1621 AGACGATCGA TGGTCAACCA AGGCGCAGAA TCAGGGAACC AGTGGAGACG AGACGCGTCG
1681 GGCAGAGATC CACTTGGACG GTTGCCCTCAG TTAGAACCGT AATCTCTTTA CGGGCAGATT
1741 GCCGTACGGG TCTTCTGCTC ATTCACTGGC AGTGCCACCG CAGTGTATC TGAGGCTGTG
1801 GCTTCAGGAT GTGGTAGCCA GTTACCATGG TCACTTGCCC TCGCTAGGAC AGCCTTCGCA
1861 GGGAAATGTC ACAGTAGCCT GCATTATGTG GTGTGAGAAC TGCTAGCGCA TTCTGTAGT
1921 TGCTTTTACG AAGCAGGATA CGCAGCGTGC ATCAGCGGGT GGTTCGAGCG CTCGCTACGC
1981 ATCACAGGGC TGTGAGGCAA GTTAGTATCT TTGGGGGACG AGTTGAGAGT GTCAGAATCG
2041 ATAGTCTCAG GGCATGCAGG CGAAATGGAG GCTGCGCCAG TAGTGCCAGC CGGTGGCGAA
2101 GCGCTCAAAT TTAATTTTTT TGTGCTGGG GATATTGTTA GAGCAACAAC TTGGGTCTAG
2161 ATGCTACTGA TAAAAAAA AA (SEQ ID NO:1).

FIG. 1

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1 CCTCGTTTTG CTTCAATCCC CGCCTTTTCT CTGTAGCTAA CCAAAGGAAC AAAGTCAGCG
61 GTAGAAGCCG TTTCTTCTGT CCGCTTCCCA CTCTTCCCGT TCGGCTGCCC CTGCAGAGCG
121 CCCTTTCTAT GCGTTGCCAC CCGTCTGCAA GTATCGCGTC TTTCGTCTCA TCAGTGATTT
181 TCTTTGCGTG TCGCGTTCGG GACGCCCTTT TCTCTCTCA ACTAACTAGC AGACGTTTCT
241 TCCGTCCCGC ATGCGACAGC GAAGGGCAGC TCCCCCAGT TCTTCATCGC CCACCTGTTG
301 TGCAACTTGT CGCCCGTCGT TCTTCACTTC TTCTCTCCCA TCCTCTCGTG ACTCTTCCTC
361 TCGAGAACTC TTTCTGTCGA ACTCTCAACC CCCACGACTG CTGGTTTCGT GGCCGTCCCCG
421 CATGCACCTT GTGTCCCGCC GCCTTGGCGC AAACACCCGC TTTCTCTGCT GTCCGCCTCC
481 CCGTGGACTT CTCTCCGTGT TTTTTCGTGT TGCCAAAAGT TTGTCTGCTT TGACGTTTCT
541 CTGCTCACC ATTGCCCGCT CTTGATGAGG AACGCTCCAC ATTGACAGCG AACTCACAGC
601 ACGCACCTC CGCGAGCGGA CTTTACAGAG CGAGGCAAGA ATCCATCGTC ACCCCGCCTA
661 CACGTACACT ACTCCACTTG GGTGCCCACG CGCGGCTTCT GGGAGACAGA GACGGTCCCTC
721 GTTTTCCGTG TCAGAACTTT GTCGAGGAAA CGCTGCTGCT GGCAGCGGGG ATTGTGACCC
781 CCCTCGGCGA ACGGGCGAAG CCGCCCTGTC GCGCGTCGCG ACTCAGCTGA GGCAGACGGC
841 GGTGCGCGGC GTGACCTCTC TTTCTTTTGT CATTCGGCCC TGATTGCAGC ACGAAGGATG
901 GAGGTCAGGG TCGGAGGCAA GTACCGACTT GGTGGAAGA TCGGCAGCGG GTCATTCGGT
961 GATATTTATA TCGGTGCAA CATTTTGACG GGGGATGAGG TGGCGATCAA GTTGGAGTCT
1021 ATCAAGTCGA AGCACCCGCA GCTGCTCTAT GAGTCGAAGC TGTACAACT GCTGGCTGGC
1081 GGCATTGGGA TTCCCATGGT CCACTGGTAC GGCATCGAAG GAGACTACAA TGTTATGGTT
1141 ATCGACCTTC TCGGCCCTTC TCTGGAGGAC CTTTTAGTA TCTGCAATCG CAAACTCTCT
1201 CTCAAGACGG TGTGATGCT CGCAGACCAG ATGCTCAACC GCATCGAGTT TGTCCATAGC
1261 AAGAACTTCA TCCATCGCGA TATCAAACCC GACAACTTCC TCATTGGCCG TGGAAAGAAG
1321 ATGTCGTCG TCTACATCAT CGATTTCCGT TTGGCAAAGA AATATCGAGA CCCAAAGACT
1381 CAGCAACATA TCCCATACAG GGAAGGCAAG AACCTAACAG GCACAGCGCG TTACGCTTCC
1441 ATCAACACCC ACCTGGGGAT CGAGCAGAGT CGGCGAGACG ACCTAGAGGC GCTCGGTTAC
1501 GTTCTCATGT ACTTCAATAG AGGTTCTCTT CCGTKGCAGG GTCTGAAGGC GACGACGAAG
1561 AAGGACAAAT ACGACAAGAT TATGGAGAAG AAAATGTCTA CTCCCATCGA AATTTTGTGC
1621 AAGCATTTCC CATTCGAGTT CATCACCTAC TTGAATTACT GCCGGTCCCT GCGCTTCGAG
1681 GATCGTCCTG ACTACGCATA CTTGCGACGC CTGTTCAAAG ACTTGTTTTT TAGAGAGGGA
1741 TATCAGTACG ACTTCATCTT CGACTGGACT TTCATCAACA CGGAGAAGGA TCGCGCGAGT
1801 CGAAGAAGCC AGCAAGTTTA TGTGGAAGAC AACCGGCAAG TTGAGGAGAA TCAGAACGAG
1861 TTGCCGATGT AGGGTGGTCG GTGTGCGGAG GCCGGCGGGG AGCGTGGAGT CCGCTGAGTC
1921 TGGAAGTCTG CAGACTGTGC TCTGGCACTC GACCCACTTG TTTGTGTTTC CCTCGACTCG
1981 CGCAGGTCGA GGAAAACAGA GACGAACAGG TTACCCAGGA GTGTTTTTGG TCAGGACGCG
2041 CGTCTCCCTC TGAGTTTCGC AAAGTTGCCC CTGGAA (SEQ ID NO:3).

FIG. 2

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1 TTAACCTCA CTAAAGGGAA CAAAAGCTGG AGCTCCACCG CGGTGGCGGC GCACCGAGGA
61 AAACGCAGCT CGTAAGAGAC AGTTCTCTCG GTGAGAAGAG CTATCCGAGA AGGACACCAT
121 GGCGCACCAT CAAGACACCC GCAACCACAC GGGGGTCGGA CCTCTTCGT CTATCCCTCT
181 GAAAGATTTG AAGATCGCCG GCGTCTGGAA AATCGGCAGA AAAATCGGAT CCGGTTCTTT
241 CGGCGACATA TACAAAGGCC TGAATTCTCA GACCGGTCAG GAGGTGGCGC TGAAGGTGCA
301 AAGCACCAAG GCGAAGCATC CGCAGTTGCT GTACGAATAC AAACCTTTTGA AGCATTTGCA
361 GGGAGGAACG GGCATTGCTC AAGTGTTCTG TTGCGAGACT GCGGGCGACC ATAACATCAT
421 GGCCATGGAG TTGCTCGGAC CTCTTTTAGA GGACGCTTTC AACTTGTGCA ATCGCACCTT
481 CTCTCTCAAA ACCATTCTTC TTCTCGCCGA CCAGTTTCTG CAACGCGTCG AGTACATCCA
541 CTCCAAGAAT TTCATTACA GAGATATCAA ACCAGATAAC TTCTTTCTCG GCGGTGCCCG
601 CAATCAAAAC ACGATCTACG TGATCGACTT CGGCCTGGCG AAGAAGTTTC GCGATCCGAA
661 AACGCACCAA CATATTCCGT ACAGAGAAAA CAAGAATCTC ACGGGAACGG CCGCTACGC
721 GTCCATCAGT GCGCATCTGG GTTCCGAGCA GAGTCGCCGA GATGACCTCG AAGCAGTCGG
781 CTACGTTCTC ATGTACTTCT GTCGAGGAGG CACGCTGCCT TGGCAGGGCA TCAAAGCGAA
841 TACCAAACAG GAGAAGTACC ACAAGATCAT GGAGAAGAAG ATGTCGACGC CCGTCGAGGT
901 GCTATGCAAG GGATATCCAA GCGAATTTGC CACATACTTG CACTACTGCC GCTCCTTGCG
961 ATTCGAGGAC CGACCGGACT ACGCCTACCT CAAGCGACTC TTTCGAGATC TCTACATCAA
1021 AGAGGGCTAC GATGACAGTG ACCGCGAATT CGACTGGACA GTGAAACTTT CGTCGCGCAG
1081 TCTCGGACCG CCAAGCAGTC GAGCGCAACA TGTTTTACTG AGTCAAGACA CCCGAACGCG
1141 AGGGAAGCGG GAGACAGATC GACCTGTGCG TCGCGGAGT GCGGACCGCG AACGAGGAAT
1201 CCATTTCAGC AACGGGAACG TGGGCAATCC TTCGATGGCA ACGAACCCCG GCGGCTGTC
1261 AGTCATGGTG CATGAACGCA CGAGTCTGGT GGATCAGGGA GACCGTGGGT CCGCGCAAAC
1321 TTCTACGCGG AAAGAAGACG CGAAGGACGG CAGATGGCCA GGAGGCAGAT TTCTTGTCT
1381 TCCACTGTTA TGTCGGCGCT CTCCGACGAA GGCTTAGATG AACTGCGGAG GCGCTCCTGT
1441 CCGCGCAGTT GGCATCTCTC TCCTTCATTG TCGTTGTTCC CCTGCAACTC GAGTCCACCC
1501 TTGACATCCT CGTCTCTCTC TTCTGTGCGG TTCTCTTTT CTCTCTCTCT CCCCCTAGC
1561 TTCGTTCTCT CCTTTCTATC CTGCTTCGGC GTCGCCCTAC TTCTCTCTCT ACTTCTCTCC
1621 CTTTTGTTTT TCTTCGCGGC GTCTCTCCTT CACTCTGTCT CCGCCTCTGA CGCCGCGCGG
1681 GAGCCGTTTC CTGCAGGCAG CTCAGGCAAT ACCTGCCTGC AGGTGCCTCT CTTTTTTGAG
1741 CGTCTCTCTT TCCTCGTCGA AACGGTCCTC ACAGCTTCCT CTCCCTGGGG ACGCCGTGGG
1801 CGTAAGTTCT TTTTTTGACG GTCCCGGTGG GCTGGCGTTG TTCGCTGCC TTCCGCGCAT
1861 GCACTCCGAG CATTTTTGCC TGGCCTGGAC TTCTCCGAGC GAGAGTTGCG GTTTGGCTTC
1921 TGCATCGTCT CCTGCGCTGC TTTCATTTCT CTAGGTTTCT GCTTGCGGCC TCCGTGTACA
1981 GAAATCGGAA GGTGAAGGCG TAGTGGCCAG AGAACGAAGC AAACGAGAGA ACCACGTTCC
2041 ACCTTGTCG CACGCATGCA TCTACGCATG CACGGTATTT AAGCCGATTT TTTGTGTATG
2101 TATATAGATG TATATATATA TGTATCTACA TGTATCTACC TATATATATG TGTGTGTGTA
2161 AGTGGAAGTG TATTTTTGCA TGTGCAGAAA GCTTTCTTTT CCGCTGGCAT GCTGGAAGAA
2221 GGGCAGGAGG CGACGATCCT GCGAGTCAGG GCGTTCCCTT GTTCCAGTG AGTTAACCGA
2281 ATTGTTTATT GATATGCGTT TGCATGCATC GACAATGGAT CCTAGACACG CCCGTTTAAA
2341 ATCAGAGGTA TTCCTAAAAA AAAAAAAAAA AAA (SEQ ID NO:5).

FIG. 3

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		1	CKI β -Nt	*	75
TgCKI β (SEQ ID NO:6)	(1)		MAHQDTRNHTGVGPSSSIPLKDL	IAGVWVIGRKIGSGSFGDIYKGLNSQTGQEVAKVETKAKHPQLLYEYK	
TgCKI α (SEQ ID NO:4)	(1)		-----MEVRVGGKYRLGRKIGSGSFGDIYIGANILTGDEVAIKLESIKSKHPQLLYESK		
EtCKI α (SEQ ID NO:2)	(1)		-----MDVRVGGKYRLGRKIGSGSFGDIYLGNTISTGDEVAIKLESVRSRHPQLLYESK		
PfCKI α (SEQ ID NO:21)	(1)		-----MEIRVANKYALGKKLGSFGDIYVAKDIVTMEFAVKLESTRSKHPQLLYESK		
LmCKI-2 (SEQ ID NO:22)	(1)		-----MNVEIRVGNRYRIGQKIGSGSFGDIYHGTNIQTGDFVAIKLEQVKTTRHPQLLYESR		
TcCKI-2 (SEQ ID NO:23)	(1)		-----MSLEIRVGNRFRLGQKIGAGSFGDIYFRGTNIQTGETVAIKLEQAKTRHPQLALEAR		
			**	*****	*****
		76			150
TgCKI β (SEQ ID NO:6)	(76)		LLKHLQGG---TGIAQVFCCTAGDENIMAMELLGPSLEDVFNLCNRTFSLKTIILLADQFLQVVEYIHSKNFIH		
TgCKI α (SEQ ID NO:4)	(55)		LYKLLAGG---IGIPMVHNYGIEGDYVNMVIDLLGPSLEDLFSICNRKLSLKTVMMLADQMLNRIEFVHSKNFIH		
EtCKI α (SEQ ID NO:2)	(55)		LYKILTGG---IGIPTLYWYGIEGDYVNMIEELLGPSLEDLFSICNRKLSLKTVMMLADQMLNRIEFVHSRHFH		
PfCKI α (SEQ ID NO:21)	(55)		LYKILGGG---IGVPKVYWYGIEGFTIMVLDLLGPSLEDLFTLCNRKPSLKTVMMLADQMLNRIEFVHSKNFIH		
LmCKI-2 (SEQ ID NO:22)	(57)		FYRILSGGGGAVGIPMMFYHGVEGEPNVMIIELLGPSLEDLFSFCGRRLSLKTTMLADQMI SRIEFVHSKSVLH		
TcCKI-2 (SEQ ID NO:23)	(57)		FYRILNAGGGVVGIPNIFLYGVEGEPNVMMVMDLLGPSLEDLFSFCDRKLSLKTTMLAEQMIARIIEFVHSKSVIH		
			*****	*****	*****
		151		CKI α -It	225
TgCKI β (SEQ ID NO:6)	(148)		RDIKPDNFILOGAGNQNTIYVIDFLGAKKFRDPKTHQHIPPYRENKNTGTARYASISAHIGSEQSRDDLEAVGY		
TgCKI α (SEQ ID NO:4)	(127)		RDIKPDNFILOGAGKMSVYVIDFLGAKKYRDPKTHQHIPPYREGKNTGTARYASINTHLGIEQSRDDLEALGY		
EtCKI α (SEQ ID NO:2)	(127)		RDIKPDNFILOGAGKMSIVFAIDFLGAKKYRDPKTHQHIPPYREGKNTGTARYASVNTHLGIEQSRDDLEALGY		
PfCKI α (SEQ ID NO:21)	(127)		RDIKPDNFILOGAGKVVLIHILDFGLAKKYRDSRSHTHIPPYREGKNTGTARYASINTHLGIEQSRDDLEALGY		
LmCKI-2 (SEQ ID NO:22)	(132)		RDIKPDNFILOGAGKGGHVVYVIDFLGAKKYRDPKTHQHIPPYREGKNTGTARYASINTHLMGVEQSRDDMEGIGY		
TcCKI-2 (SEQ ID NO:23)	(132)		RDMKPDNFILOGAGKGGHVVYVIDFLGAKKYRDPKTHQHIPPYREGKNTGTARYASINTHLGIEQSRDDLEBEGY		
			*****	*****	*****
		226			300
TgCKI β (SEQ ID NO:6)	(223)		VLMYFCRGGLPWQGIKANTKQEKYHKIMEKKMSTPVEVLCKGYPSEFATYLYHCRSLRFEDRPDYAYLKRFLPD		
TgCKI α (SEQ ID NO:4)	(202)		VLMYFNRG-SLPWQGLKATTKKDKYDKIMEKKMSTPIEILCKHPPFEFITYLNYCRSLRFEDRPDYAYLRRLPKD		
EtCKI α (SEQ ID NO:2)	(202)		VLMYFNRG-SLPWQGLKATTKKDKYDKIMEKKMSTPIEVLCKQPPFEFITYLNYCRSLRFEDRPDYAYLRRLPKD		
PfCKI α (SEQ ID NO:21)	(202)		VLMYFLRG-SLPWQGLKAISKDKYDKIMEKKISTSEVLCKRNASPEFVYLYNCRSLRFEDRPDYAYLRRLLKD		
LmCKI-2 (SEQ ID NO:22)	(207)		ILMYFLRG-SLPWQGLKAHTKQEKYNRISERKQTPVELLCKGPPSEFAAYMNYVRALRFEDKPDYSYLKRMFRD		
TcCKI-2 (SEQ ID NO:23)	(207)		ILMYFLRG-SLPWQGLKAHTKQEKYSRISERKQTPVETLCKGPPAEFAAYLNYIRSLRFEDKPDYSYLKRLFRE		
			*****	*****	*****
		301		CKI α -Ct	375
TgCKI β (SEQ ID NO:6)	(298)		LYIKEGYDDSDREFDWTIKLSSRLGPPSSRAQHVLISQOTRTRGKRETDPRVAARSGDRERGIHPSNGNVGNPS		
TgCKI α (SEQ ID NO:4)	(276)		LFIREGYQY-DLFDWTFINTEKDRASRSQQVYVEDNRQVEENQNELEPM		
EtCKI α (SEQ ID NO:2)	(276)		LFIREGYQY-DLFDWTFIHAERERERQRSMVNGAEGSNQWRDASGRDPLGRLPQLEP		
PfCKI α (SEQ ID NO:21)	(276)		LFIREGYTY-DLFDWTFVYASEKDKKMLNKNRFDQTADQEGRDQRNN		
LmCKI-2 (SEQ ID NO:22)	(281)		LFVREGYHV-DYVFDWTLKRIHESLQEQSPFGSGNGGAAGNSPVNQSPAQGGNGGAPNSANNQESGAPEQQ		
TcCKI-2 (SEQ ID NO:23)	(281)		LFIREGYHV-DYVFDWTLKRIHESLQEQSPFGSGNGGAGNSPVNQSPAQGGNGGAPNSANNQESGAPEQQ		
			*****	*****	*****
		376		CKI β -Ct	435
TgCKI β (SEQ ID NO:6)	(373)		MATNPGGLSVMVHERTSLVDQDGRGRETSTRKEDAKDGRWPGGRFSCLFLICRRSPKA		

* Brackets mark the catalytic core region

FIG. 4

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	TgCKI β	TgCKI α	EtCKI α	PfCKI α	LmCKI-2	TcCKI-2
TgCKI β	100	48	48	45	44	42
TgCKI α		100	81	68	58	62
EtCKI α			100	67	59	60
PfCKI α				100	52	55
LmCKI-2					100	75
TcCKI-2						100

FIG. 5

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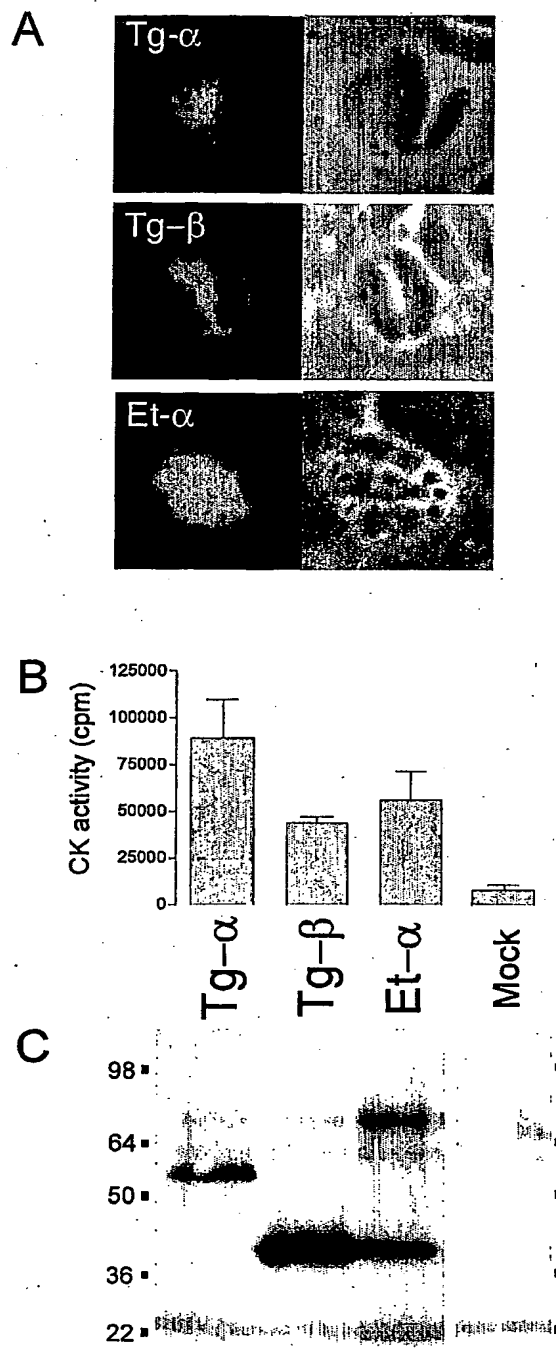


FIG. 6A-C

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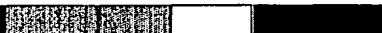
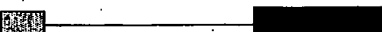

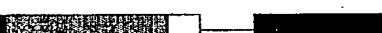
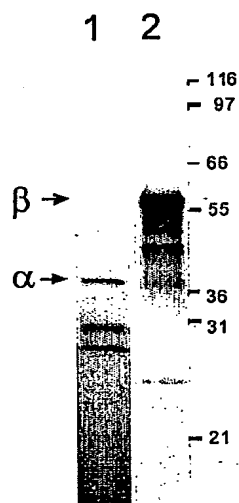
	Location	CK Activity
TgCKI α FLAG 	C	+++
TgCKI β FLAG 	M	++
TgCKI β -CAT 	M	+++
3' Δ 332-CAT 	C	-
3' Δ 182-CAT 	C	-
3' Δ 64-CAT 	C	+++

FIG. 7

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A.



B.

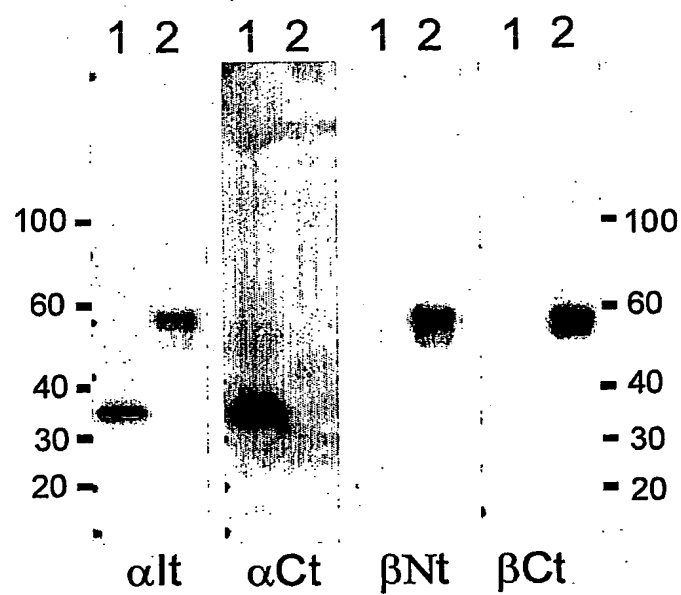


FIG. 8A-B

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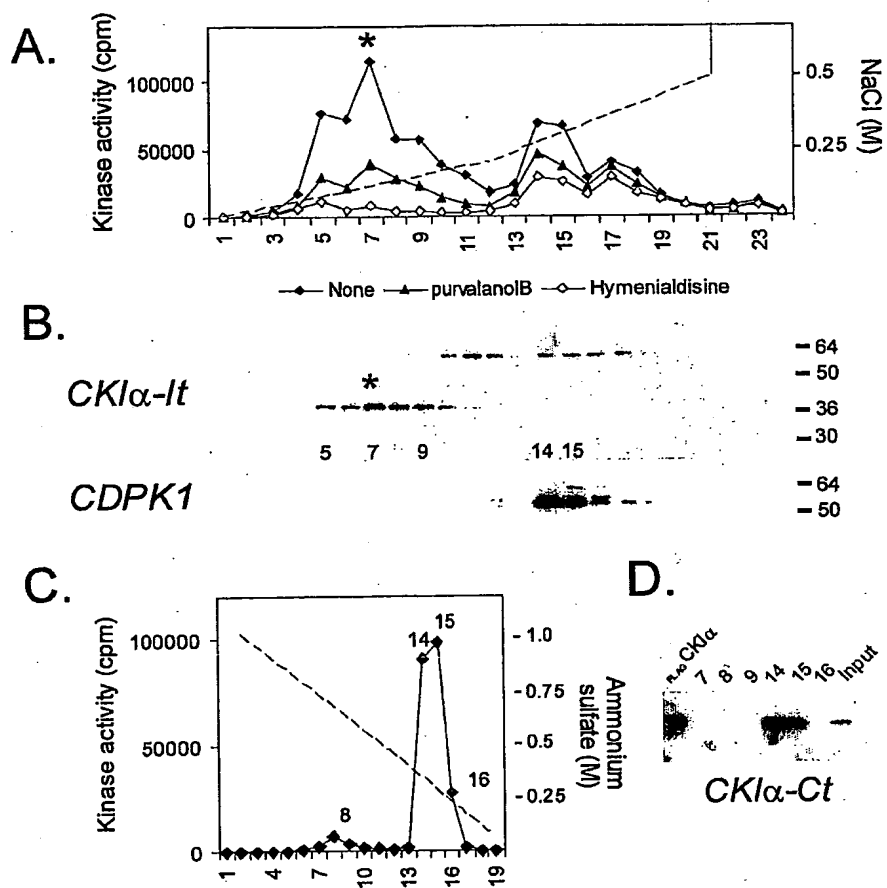


FIG. 9A-D

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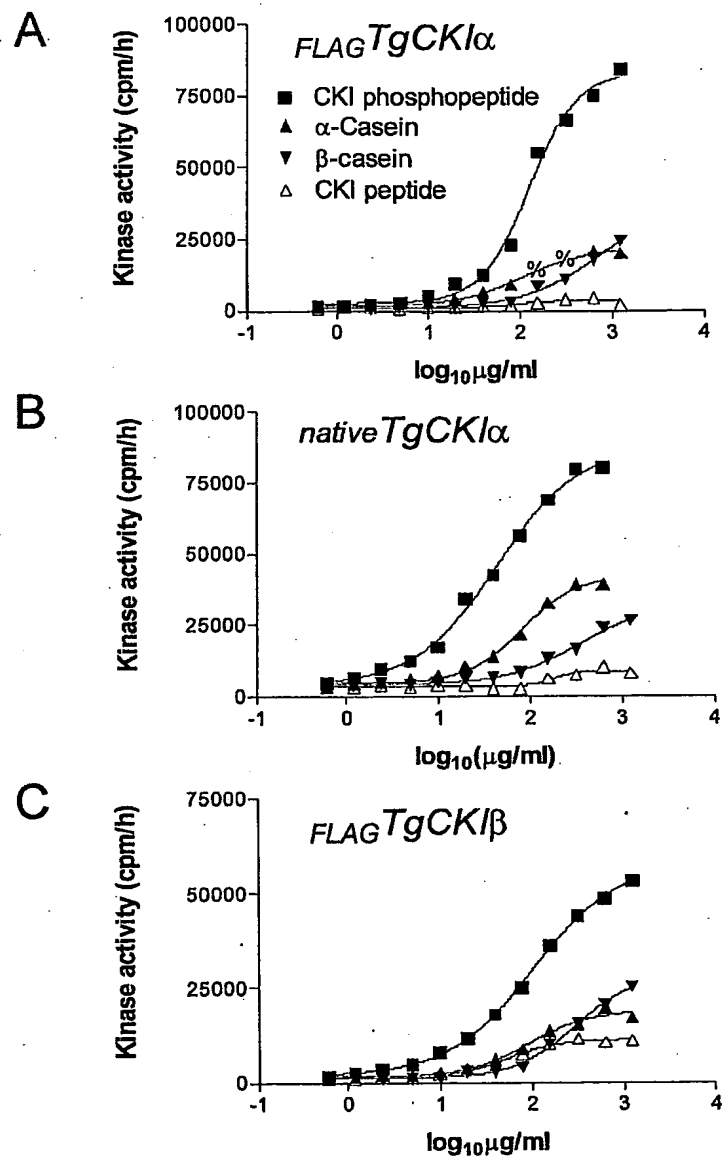
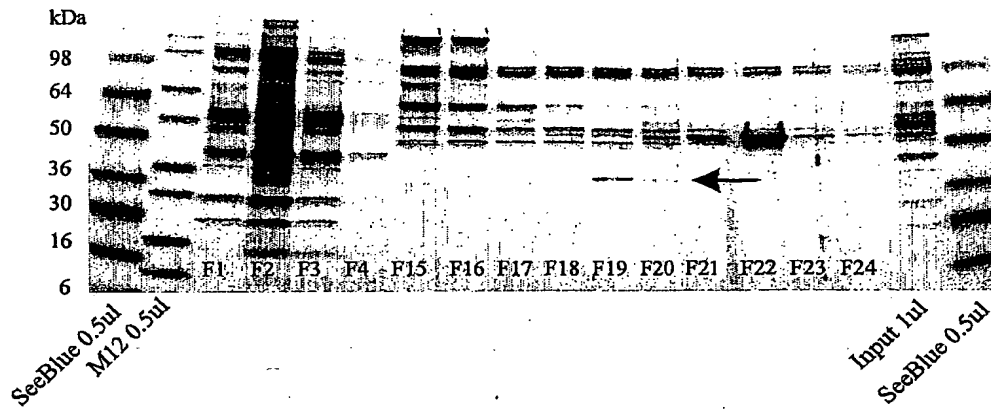


FIG. 10A-C

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A.



B.

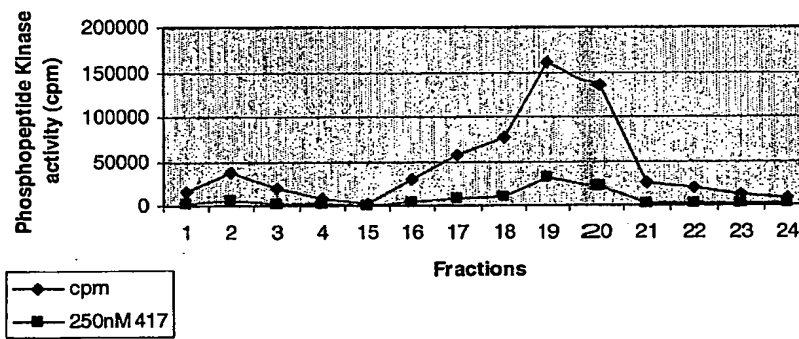


FIG. 11A-B

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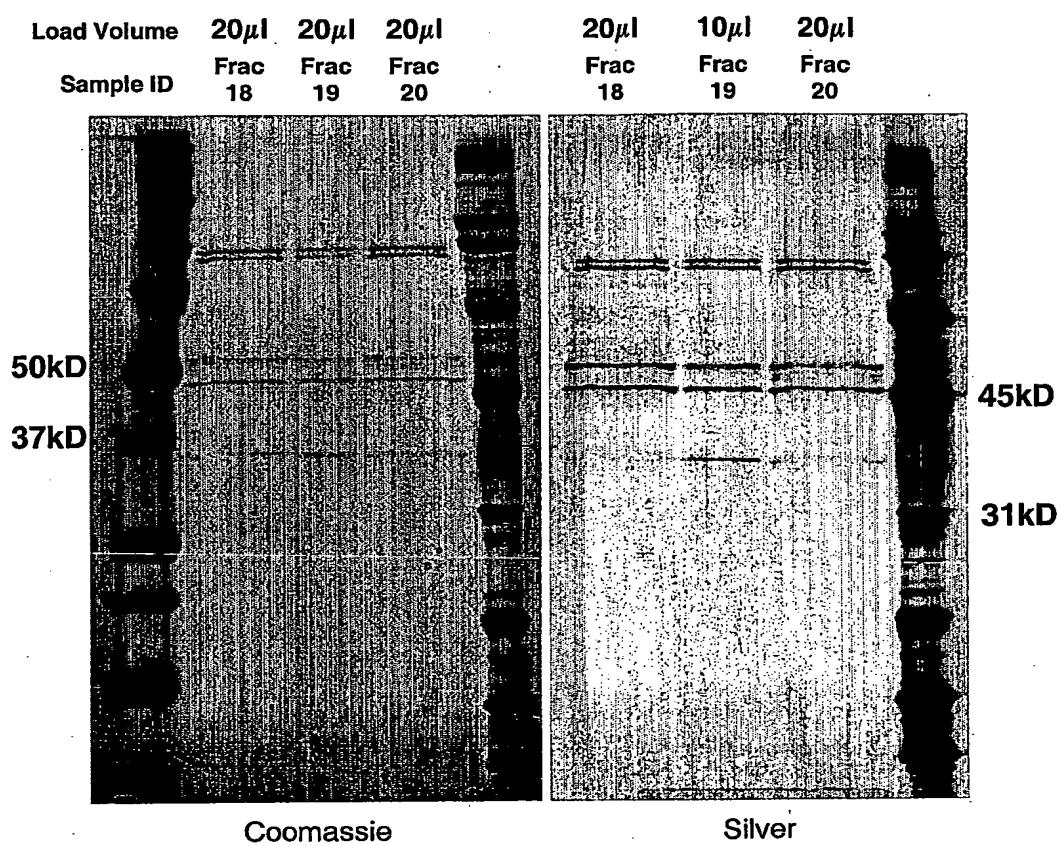


FIG. 12

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MDVRVGGKYR LGRKIGSGSF GDIYLG TNIS TGDEVAIKLE
SVRSRHPOLI YESKLYKILT GGIGIPTLYW YGIEGDYNVM
IIELLGPSLE DLFSICNRKL SLKTVLMLAD QMLNR IEFVH
SRHFIHRDIK PDNFLIGRGK KMSIVFAIDF GLAKKYRDPR
TQSHIPYREG KNLTGTARYA SVNTHLGIEQ SRRDDLEALG
YVLMYFNREG LPWQGLKATT KKDKYDKIME KKMSTPIEVL
CKQFPFEFIT YLNYCRSLRF EDRPDYSYLR RLFKDLFFRE
GYQYDFIFDW TFLHAERERE RQRSMVNOG AESGNQWRRD
ASGRDPLGRL PQLEP (SEQ ID NO:2)

FIG. 13

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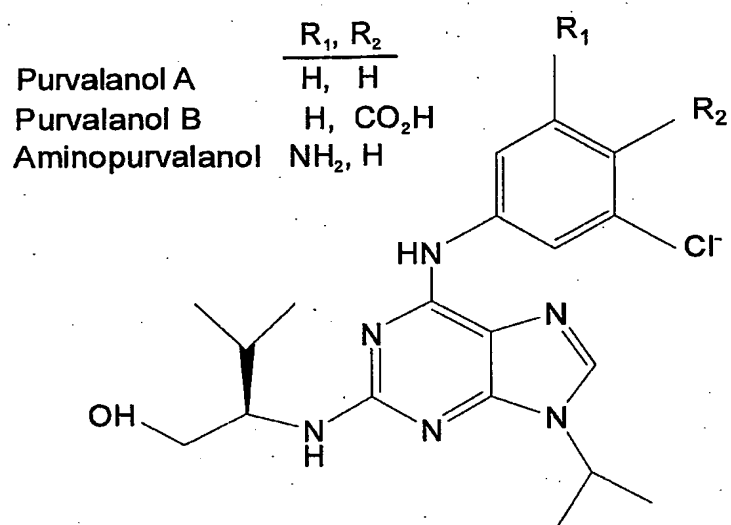


FIG. 14

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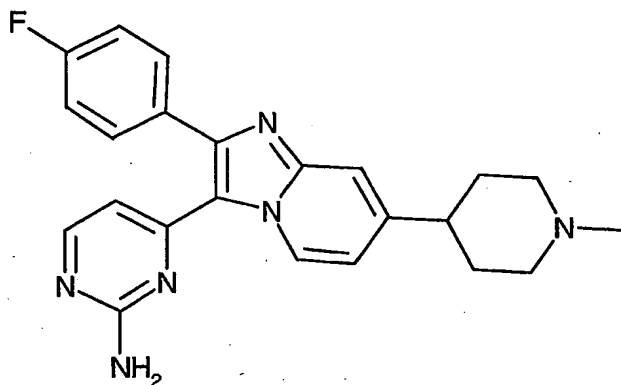


Fig. 15